

Result No.	Score	Query		Length	DB	ID	Description
		Match	%				
1	348	100.0	1637	4	US-09-949-016-2339		Sequence 2339, Ap
2	348	100.0	1642	4	US-09-159-277A-1		Sequence 1, Appli
3	348	100.0	1642	4	US-09-023-655-1485		Sequence 1485, Ap
4	348	100.0	1642	4	US-08-844-691A-1		Sequence 1, Appli
5	348	100.0	1701	3	US-09-357-072-1		Sequence 1, Appli
6	348	100.0	1701	3	US-08-983-502-1		Sequence 1, Appli
7	348	100.0	1701	4	US-09-516-747-1		Sequence 1, Appli
8	348	100.0	1701	4	US-09-933-814-1		Sequence 1, Appli
9	348	100.0	1701	5	PCT-US96-16542-1		Sequence 1, Appli
10	348	100.0	1701	5	PCT-US96-10521-1		Sequence 1, Appli
11	346.4	99.5	606	3	US-09-064-414-1		Sequence 1, Appli
12	344.8	99.1	606	3	US-09-064-414-3		Sequence 3, Appli
13	344.8	99.1	627	3	US-09-064-414-5		Sequence 5, Appli
14	307	88.2	8025	1	US-09-949-016-14081		Sequence 14081, A
15	43.4	12.5	7218	1	US-08-232-463-14		Sequence 14, Appli
16	43.2	12.4	4284	4	US-09-502-540-3289		Sequence 3289, Ap
17	43.2	12.4	17727	4	US-09-502-540-1152		Sequence 1152, Ap
18	38.2	11.0	801	4	US-08-252-991A-5148		Sequence 5148, Ap
19	38.2	11.0	1269	4	US-09-252-991A-5037		Sequence 5037, Ap
20	38.2	11.0	2031	4	US-09-252-991A-5180		Sequence 5180, Ap
21	36.6	10.5	4058	4	US-09-774-528-135		Sequence 135, App
22	36.2	10.4	596	3	US-09-385-982-304		Sequence 304, App
23	35.2	10.1	939	4	US-09-489-039A-2942		Sequence 2942, Ap
24	35	10.1	601	4	US-09-949-016-187647		Sequence 187647,
25	35	10.1	2751	3	US-09-037-190-45		Sequence 45, Appli
26	35	10.1	2751	3	US-09-037-192-45		Sequence 45, Appli
27	35	10.1	2751	3	US-09-037-143-45		Sequence 45, Appli

Db  
668 CAGGCCCGTGACTCTCCAGAACAGGAGTGGGGCCATGTCCCCGATGTCA 715

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RESULT 2
US-09-159-277A-1
; Sequence 1, Application US/09159277A
; Patent No. 6562797
; GENERAL INFORMATION:
; APPLICANT: DIXIT, VISHVA M.
; APPLICANT: O'ROURKE, KAREN
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR REGULATING
; TITLE OF INVENTION: FAS-ASSOCIATED APOPTOSIS
; NUMBER OF SEQUENCES: 10
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Morrison & Foerster LLP
; STREET: 755 Page Mill Road
; CITY: Palo Alto
; STATE: CA
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patent In Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/159,277A
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/844,691
; FILING DATE: 21-APR-1997
; APPLICATION NUMBER: US 08/416,379
; FILING DATE: 03-APR-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Konski, Antoinette F.
; REGISTRATION NUMBER: 34,202
; REFERENCE/DOCKET NUMBER: 203442107001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (650)813-5600
; TELEFAX: (650)494-0792
; TELEX: 706141 MRSNFOERS SFO
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1642 base pairs
; TYPE: nucleic acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: cDNA
; FEATURE:
; NAME/KEY: CDS
; LOCATION: 130..753
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 4..6
; OTHER INFORMATION: /note= "An in-frame stop codon 130
; OTHER INFORMATION: base pairs upstream of the initiator me
; FEATURE:
; NAME/KEY: polyA signal
; LOCATION: 1636..1641
; OTHER INFORMATION: /note= "Potential poly(A) adenylation s
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 198..753
; OTHER INFORMATION: /note= "Clone-15; 5' end of FADD"
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 249..753
; OTHER INFORMATION: /note= "Clone-8; 5' end of FADD"
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 460..660

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; OTHER INFORMATION: /note= "Death Domain of FADD"
; FEATURE:
; NAME/KEY: mutation
; LOCATION: replace(490..492, "aay")
; OTHER INFORMATION: /note= "For FADDmt: sequence is altered to either AAT or
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: group(250..753, 232..753)
; OTHER INFORMATION: /note= "Codons can comprise C-terminal polypeptide fragme
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: 460..639
; OTHER INFORMATION: /note= "death domain homology region"
US-09-159-277A-1

Query Match      100.0%; Score 348; DB 4; Length 1642;
Best Local Similarity 100.0%; Pred. NO. 3.8e-94;
Matches 348; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY    1   TTGAGCGGGGGCGGGCGGGCGGGCGCCCTGGGGGAAGAAGACCTGTGTGCAGCATTTT 60
Db     373  TTGAGCGGGGGCGGGCGGGCGGGCGCCCTGGGGGAAGAAGACCTGTGTGCAGCATTT 432

QY    61   AACCTCATATGTGTAATGTGGGAAAGATTGAGAAAGCTGGCTCCTCAGCTCAAAAGTC 120
Db     433  AACGTCTATATGTGTAATGTGGGAAAAGATTGAGAAAGCTGGCTCCTCAGCTCAAAGTC 492

QY    121  TCAGACACCAGATCGACAGCATCGAGGACAGATATCCCCGCCAACCTGCAGAGCGGTGTG 180
Db     493  TCAGACACCAGATCGACAGCATCGAGGACAGATATCCCCGCCAACCTGCAGAGCGGTGTG 552

QY    181  CGGGAGTCACTGAGAAATCTGGAAGAACACAGAGAAGAGAACGCAACAGTGCGCCACTTG 240
Db     553  CGGGAGTCACTGAGAAATCTGGAAGAACACAGAGAAGAGAACGCAACAGTGCGCCACTTG 612

QY    241  GTGGGGCTCTCAGGTCTGCCAGATGAACCTGGTGGCTGACCTGGTACAAGAGGTTTCAG 300
Db     613  GTGGGGCTCTCAGGTCTGCCAGATGAACCTGGTGGCTGACCTGGTACAAGAGGTTTCAG 672

QY    301  CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCAGATGTCA 348
Db     673  CAGGCCCGTGACCTCCAGAACAGGAGTGGGGCCATGTCCCAGATGTCA 720


RESULT 3
US-09-023-655-1485
; Sequence 1485, Application US/09023655
; Patent No. 6607879
; GENERAL INFORMATION:
; APPLICANT: Cocks, Benjamin G.
; APPLICANT: Susan G. Stuart
; APPLICANT: Jeffrey J. Seilhamer
; TITLE OF INVENTION: COMPOSITION FOR THE DETECTION OF BLOOD CELL GENE
; , TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 1508
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: INCYTE PHARMACEUTICALS, INC.
; STREET: 3174 PORTER DRIVE
; CITY: PALO ALTO
; STATE: CALIFORNIA
; COUNTRY: USA
; ZIP: 94304
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Word Perfect 6.1 for Windows/MS-DOS 6.2
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/023,655
; FILING DATE: HERewith
; CLASSIFICATION:
; PRIOR APPLICATION NUMBER:
; APPLICATION NUMBER:

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	Query Match	100.0%	Score 348;	DB 4;	Length 1642;
	Best Local Similarity	100.0%;	Pred. No. 3.8e-94;		
	Matches 348;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0
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Dd	373				
		TTCAGGCGGGGGCGGGCGGGCGGGCGGGCTGTGGGAAGAAGACTGTGTGCAGCATTT	432		
Qy	61	AACGTCATATGTGATAATGTGGGGAAAGATTGGAGAAGGCTGGCTCAGCTCAAAGTCT	120		
Dd	433	AACGTCATATGTGATAATGTGGGGAAAGATTGGAGAAGGCTGGCTCAGCTCAAAGTCT	492		
Qy	121	TCAGACACCAAGATCGACAGCATCGAGACAGATACCCCGAACCTGACACAGCGGTGTG	180		
Dd	493	TCAGACACCAAGATCGACAGCATCGAGACAGATACCCCGAACCTGACAGCGGTGTG	552		

	Query Match	100.0%	Score 348;	DB 4;	Length 1642;
	Best Local Similarity	100.0%;	Pred. No. 3.8e-94;		
	Matches 348;	Conservative 0;	Mismatches 0;	Indels 0;	Gaps 0;
Qy	1	TTGAGCGGGGGCGGGCGGGCCGCCGCTGGGGGAAGAAGACCTGTGTGCAGCATTT	60		
Dd	373	TTGAGCGGGGGCGGGCGGGCCGCCGCTGGGGGAAGAAGACCTGTGTGCAGCATTT	432		
Qy	61	AACGTCATATGTGATAATTGGGGAAGAAGATTGGAAAGCTGGCTCGTCAGCTCAAAGTC	120		
Dd	433	AACGTCATATGTGATAATTGGGGAAGAAGATTGGAAAGCTGGCTCGTCAGCTCAAAGTC	492		
Qy	121	TCAGACCAAGATCGACAGCATCGAGACAGATACCCTCCGAACTGACACAGCGTGTG	180		
Dd	493	TCAGACCAAGATCGACAGCATCGAGACAGATACCCTCCGAACTGACAGCGTGTG	552		















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